

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/744,804

CRF Processing Date: 1/15/2002
 Edited by: _____
 Verified by: [Signature] (STIC staff)

ENTERED

PU109

JA

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was wrapped down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____.
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

PCT09

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/744,804

DATE: 01/15/2002

TIME: 13:01:47

Input Set : A:\EISENBACH3.txt

Output Set: N:\CRF3\01152002\I744804.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Yeda Research & Development Company Limited at the
 4 Bio-Technology General Corp.
 6 <120> TITLE OF INVENTION: Tumor Associated Antigen Peptides and Use of Same as
 7 Anti-Tumor Vaccines
 9 <130> FILE REFERENCE: 453-A-PCT
 11 <140> CURRENT APPLICATION NUMBER: US/09/744,804
 12 <141> CURRENT FILING DATE: 2001-01-30
 14 <150> PRIOR APPLICATION NUMBER: 125608
 15 <151> PRIOR FILING DATE: 1998-07-30
 17 <160> NUMBER OF SEQ ID NOS: 77
 19 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

857 <210> SEQ ID NO: 77
 858 <211> LENGTH: 9
 859 <212> TYPE: PRT
 860 <213> ORGANISM: Homo sapiens
 862 <400> SEQUENCE: 77
 863 Met Leu Ala Gly Ile Cys Leu Ser Ile
 864 5
 E--> 871 15

VERIFICATION SUMMARY

DATE: 01/15/2002

PATENT APPLICATION: US/09/744,804

TIME: 13:01:48

Input Set : A:\EISENBACH3.txt

Output Set: N:\CRF3\01152002\I744804.raw

L:11 M:270 C: Current Application Number differs, Replaced Application Number

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:871 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:77